

CRF Errors Corrected by the STIC Systems Branch

 CRF Processing Date: 4/3/03
 Edited by: DC
 Verified by: DC (STIC staff)
Serial Number: 10/030,014

- ☐ Changed a file from non-ASCII to ASCII
- ☐ Changed the margins in cases where the sequence text was "wrapped" down to the next line.
- ☐ Edited a format error in the Current Application Data section, specifically: _____
- ☐ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other _____
- ☐ Added the mandatory heading and subheadings for "Current Application Data".
- ☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
- ☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically: _____
- ☐ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: _____
- ☐ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: _____
- ☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
- ☐ Inserted colons after headings/subheadings. Headings edited included: _____
- ☐ Deleted extra, invalid, headings used by an applicant, specifically: _____
- ☐ Deleted: ☐ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file; ☐ page numbers throughout text; ☐ other invalid text, such as _____
- ☐ Inserted mandatory headings, specifically: _____
- ☐ Corrected an obvious error in the response, specifically: _____
- ☐ Edited identifiers where upper case is used but lower case is required, or vice versa.
- ☐ Corrected an error in the Number of Sequences field, specifically: _____
- ☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
- ☐ Deleted **ending** stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: _____
- ☒ Other: Re-aligned. amino numbering in Seq. 538

*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.



PCT10

RAW SEQUENCE LISTING

DATE: 04/03/2003

PATENT APPLICATION: US/10/030,014

TIME: 11:24:28

Input Set : N:\Darika\10030014DC.txt

Output Set: N:\CRF4\04032003\J030014.raw

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3 <110> APPLICANT: Epimmune Inc.
4   Sette, Alessandro
5   Sidney, John
6   Southwood, Scott
9 <120> TITLE OF INVENTION: HLA Binding Peptides and Their Uses
12 <130> FILE REFERENCE: 39963-20021.00
14 <140> CURRENT APPLICATION NUMBER: US 10/030,014
15 <141> CURRENT FILING DATE: 2001-12-28
17 <150> PRIOR APPLICATION NUMBER: PCT/US00/17842
18 <151> PRIOR FILING DATE: 2000-06-28
20 <150> PRIOR APPLICATION NUMBER: US 60/141,422
21 <151> PRIOR FILING DATE: 1999-06-29
23 <160> NUMBER OF SEQ ID NOS: 538
25 <170> SOFTWARE: FastSEQ for Windows Version 3.0
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29 <212> TYPE: PRT
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32 <220> FEATURE:
33 <223> OTHER INFORMATION: HBV.core.18.Q2Y6, peptide 1302.01
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37   1             5             10
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44 <220> FEATURE:
45 <223> OTHER INFORMATION: HBV.env.183.Y1, peptide 1302.02
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49   1             5
51 <210> SEQ ID NO: 3
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56 <220> FEATURE:
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61   1             5
63 <210> SEQ ID NO: 4
64 <211> LENGTH: 9

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85 1 5
87 <210> SEQ ID NO: 6
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90 <213> ORGANISM: Artificial Sequence
92 <220> FEATURE:
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111 <210> SEQ ID NO: 8
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121 1 5
123 <210> SEQ ID NO: 9
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125 <212> TYPE: PRT
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128 <220> FEATURE:
129 <223> OTHER INFORMATION: HCV.NS3.1078, peptide F158.01

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176 <220> FEATURE:
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184 <211> LENGTH: 9
185 <212> TYPE: PRT
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189 <223> OTHER INFORMATION: mp53.7.V9, peptide 1349.05
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192 Ser Gln Ser Asp Ile Ser Leu Glu Val
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237 <223> OTHER INFORMATION: mp53.100.V9, peptide 1349.11
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255 <210> SEQ ID NO: 20
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257 <212> TYPE: PRT
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Input Set : N:\Darika\10030014DC.txt

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261 <223> OTHER INFORMATION: mp53.119, peptide 1349.13
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267 <210> SEQ ID NO: 21
268 <211> LENGTH: 9
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276 Val Met Cys Thr Tyr Ser Pro Pro Val
277 1 5
279 <210> SEQ ID NO: 22
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282 <213> ORGANISM: Artificial Sequence
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294 <213> ORGANISM: Artificial Sequence
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300 Lys Leu Phe Cys Gln Leu Ala Lys Thr
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VERIFICATION SUMMARY

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